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## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	10/649,952
Source:	01/4
Date Processed by STIC:	.9/5/2003
•	

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<a href="http://www.uspto.gov/ebc/efs/downloads/documents.htm">http://www.uspto.gov/ebc/efs/downloads/documents.htm</a>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- Hand Carry directly to:
   U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7<sup>th</sup> Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
  - U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
- 4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 04/24/2003



OIPE

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DATE: 09/05/2003
                      RAW SEQUENCE LISTING
                      PATENT APPLICATION: US/10/649,952
                                                                TIME: 08:59:55
                      Input Set : A:\PTO.YF.txt
                      Output Set: N:\CRF4\09052003\J649952.raw
      3 <110> APPLICANT: SUNTORY LIMITED
              SUNTORY BIOMEDICAL RESEARCH LIMITED
      6 <120> TITLE OF INVENTION: Promoters of the growth and/or differentiation of
hematopoietic stem
              cells and/or
              hematopoietic progenitors
      8
     10 <130> FILE REFERENCE: YCT-785
    12 <140> CURRENT APPLICATION NUMBER: US/10/649,952

★ 12 <141> CURRENT FILING DATE: 2003-08-29

     12 <150> PRIOR APPLICATION NUMBER: JP 2001-400330
W--> 13 <151> PRIOR APPLICATION NUMBER: JP 2001-400330

W--> 13 <151> PRIOR FILING DATE: December 28, 2001 2001-12-28 ( use this date format
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       (2107
ERRORED SEQUENCES
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     18 <211> LENGTH: 166
                                                                Corrected Diskette Needec
     19 <212> TYPE: PRT
                                                                             mandatory

nunevie

identifier

wherever <2217,

12227, or <2237

is shown
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     21 <223> OTHER INFORMATION: Amino acid sequence of human Cofilin.
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     26 Asp Met Lys Val Arg Lys Ser Ser Thr Pro Glu Glu Val Lys Lys Arg
                                                                30
                                           25
     28 Lys Lys Ala Val Leu Phe Cys Leu Ser Glu Asp Lys Lys Asn Ile Ile
     29
                  35
     30 Leu Glu Glu Gly Lys Glu Ile Leu Val Gly Asp Val Gly Gln Thr Val
     31
                                   55
     32 Asp Asp Pro Tyr Ala Thr Phe Val Lys Met Leu Pro Asp Lys Asp Cys
                                                                         80
                                                    75
                              70
     34 Arg Tyr Ala Leu Tyr Asp Ala Thr Tyr Glu Thr Lys Glu Ser Lys Lys
                                                90
     36 Glu Asp Leu Val Phe Ile Phe Trp Ala Pro Glu Ser Ala Pro Leu Lys
                                          105
                     100
     38 Ser Lys Met Ile Tyr Ala Ser Ser Lys Asp Ala Ile Lys Lys Leu
                                      120
                                                           125
                 115
     40 Thr Gly Ile Lys His Glu Leu Gln Ala Asn Cys Tyr Glu Glu Val Lys
                                                       140
                                  135
     42 Asp Arg Cys Thr Leu Ala Glu Lys Leu Gly Gly Ser Ala Val Ile Ser
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44 Leu Glu Gly Lys Pro Leu

DATE: 09/05/2003

TIME: 08:59:55

26

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Input Set : A:\PTO.YF.txt
                    Output Set: N:\CRF4\09052003\J649952.raw
    45 (2107
                        165
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     48 <211> LENGTH: 501
     49 <212> TYPE: DNA
     51 <223> OTHER INFORMATION: Nucleotide sequence of human Cofilin.
€ > 53 <400> SEQUENCE: 2
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     55 cgtaagtett caacgecaga ggaggtgaag aagegeaaga aggeggtget ettetgeetg 120
     56 agtgaggaca agaagaacat catcctggag gagggcaagg agatcctggt gggcgatgtg 180
     57 ggccagactg tcgacgatcc ctacgccacc tttgtcaaga tgctgccaga taaggactgc 240
     58 cgctatgccc tctatgatgc aacctatgag accaaggaga gcaagaagga ggatctggtg 300
     59 tttatcttct gggcccccga gtctgcgccc cttaagagca aaatgattta tgccagctcc 360
     60 aaggacgcca tcaagaagaa gctgacaggg atcaagcatg aattgcaagc aaactgctac 420
     61 gaggaggtca aggaccgctg caccctggca gagaagctgg ggggcagtgc ggtcatctcc 480
                                                                         501
     62 ctggagggca agcctttgtg a
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     66 <212> TYPE: DNA
     67 <213> ORGANISM: Artificial sequence
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     68 223> OTHER INFORMATION: Primer sequence for amplifying human Cofilin gene.
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E--> 73 (200)/4 /2/0
     74 <211> LENGTH: 26
     75 <212> TYPE: DNA
     76 <213> ORGANISM: Artificial sequence
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/649,952

**€**> 79 <400> SEQUENCE: 4

80 tctccctgga gggcaagcct ttgtga

DATE: 09/05/2003

## VERIFICATION SUMMARY

PATENT APPLICATION: US/10/649,952 TIME: 08:59:56

Input Set : A:\PTO.YF.txt

Output Set: N:\CRF4\09052003\J649952.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:13 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD L:17 M:250 E: Invalid Numeric Identifier, INVALID IDENTIFIER L:23 M:282 E: Numeric Field Identifier Missing, <210> is required. L:23 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:0 L:23 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:0 differs:1 L:47 M:250 E: Invalid Numeric Identifier, INVALID IDENTIFIER L:53 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:0 L:53 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:0 differs:2 L:64 M:250 E: Invalid Numeric Identifier, INVALID IDENTIFIER L:68 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:0 L:70 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:0 L:70 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:0 differs:3 L:71 M:254 E: No. of Bases conflict, LENGTH:Input:25 Counted:26 SEQ:3 L:71 M:252 E: No. of Seq. differs, <211> LENGTH:Input:25 Found:26 SEQ:3 L:73 M:250 E: Invalid Numeric Identifier, INVALID IDENTIFIER L:77 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:0

L:79 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:0 differs:4

L:79 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:0